RAW SEQUENCE LISTING PATENT APPLICATION US/08/908,867A

DATE: 05/03/1999 TIME: 17:36:56

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This Raw Listing contains the General Information Section and up to the first 5 pages.

```
SEQUENCE LISTING
 1
 2
 3
    (1)
            General Information:
         (i) APPLICANT: YOUNG, Andrew A.
 6
                        GEDULIN, Bronislava
 7
                        BEELEY, Nigel Robert Arnold
                                                                 ENTERED
 8
                        PRICKETT, Kathryn S.
 9
        (ii) TITLE OF INVENTION: METHODS FOR REGULATING
10
                                  GASTROINTESTINAL MOTILITY
11
12
        (iii) NUMBER OF SEQUENCES: 37
13
         (iv) CORRESPONDENCE ADDRESS:
15
               (A) ADDRESSEE: LYON & LYON
16
               (B) STREET: 633 WEST FIFTH STREET
17
18
               (C) CITY: LOS ANGELES
19
               (D) STATE: CALIFORNIA
20
               (E) COUNTRY: USA
21
               (F) ZIP: 90017
22
23
          (v) COMPUTER READABLE FORM:
24
               (A) MEDIUM TYPE: Floppy disk
25
               (B) COMPUTER: IBM PC compatible
26
               (C) OPERATING SYSTEM: PC-DOS/MS-DOS
               (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
27
28
        (vi) CURRENT APPLICATION DATA:
29
               (A) APPLICATION NUMBER:
30
                                        08/908,867
               (B) FILING DATE: 08-AUGUST-1997
               (C) CLASSIFICATION: Pending
       (vii) PRIOR APPLICATION DATA:
               (A) APPLICATION NUMBER: 08/694,954
               (B) FILING DATE: 08-AUGUST-1996
36
               (C) CLASSIFICATION: Pending
      (viii) ATTORNEY/AGENT INFORMATION:
39
               (A) NAME: DUFT, BRADFORD J.
40
               (B) REGISTRATION NUMBER: 32,219
41
               (C) REFERENCE/DOCKET NUMBER: 227/166
42
43
        (ix) TELECOMMUNICATION INFORMATION:
44
45
               (A) TELEPHONE: 619/552-2200
               (B) TELEFAX: 213/955-0440
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47
               (C) TELEX: 67-3510
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     (2) INFORMATION FOR SEQ ID NO:1:
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51
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               (A) LENGTH: 39 amino acids
53
               (B) TYPE: amino acid
54
55
               (C) STRANDEDNESS: single
56
               (D) TOPOLOGY: linear
57
          (ii) MOLECULE TYPE: peptide
58
59
          (ix) FEATURE:
60
61
               (B) LOCATION: 39
               (D) OTHER INFORMATION: amidated Ser (Serineamide)
62
63
64
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
65
    His Ser Asp Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
66
67
                                          10
    Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
68
69
                 20
                                      25
    Ser Gly Ala Pro Pro Pro Ser
70
71
             35
72
73
    (2) INFORMATION FOR SEQ ID NO:2:
74
75
76
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               (A) LENGTH: 39 amino acids
77
               (B) TYPE: amino acid
78
               (C) STRANDEDNESS: single
79
80
               (D) TOPOLOGY: linear
81
82
         (ii) MOLECULE TYPE: peptide
83
         (ix) FEATURE:
84
               (B) LOCATION: 39
85
86
               (D) OTHER INFORMATION: amidated Ser (Serineamide)
87
88
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
89
    His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
90
91
                                          10
    Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
92
93
                 20
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                                                            30
94
    Ser Gly Ala Pro Pro Pro Ser
95
             35
96
97
    (2) INFORMATION FOR SEQ ID NO:3:
98
99
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(i) SEQUENCE CHARACTERISTICS:
100
                (A) LENGTH: 30 amino acids
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                (B) TYPE: amino acid
102
103
                (C) STRANDEDNESS: single
104
                (D) TOPOLOGY: linear
105
106
         (ii) MOLECULE TYPE: peptide
107
108
         (ix) FEATURE:
109
                (B) LOCATION: 30
                (D) OTHER INFORMATION: amidated Arg (Arginineamide)
110
111
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
112
113
     His Ala Glu Gly Thr Phe Thr Ser Asp Val Ser Ser Tyr Leu Glu Gly
114
115
                                           10
     Gln Ala Ala Lys Glu Phe Ile Ala Trp Leu Val Lys Gly Arg
116
                                     25
117
118
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     (2) INFORMATION FOR SEQ ID NO:4:
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121
122
           (i) SEQUENCE CHARACTERISTICS:
                (A) LENGTH: 31 amino acids
123
                (B) TYPE: amino acid
124
                (C) STRANDEDNESS: single
125
                (D) TOPOLOGY: linear
126
127
         (ii) MOLECULE TYPE: peptide
128
129
         (ix) FEATURE:
130
131
                (B) LOCATION: 31
132
                (D) OTHER INFORMATION: amidated Ser (Serineamide)
133
134
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
135
     Asp Leu Ser Lys Gly Met Glu Glu Glu Ala Val Arg Leu Phe Ile Glu
136
137
                                          10
138
     Trp Leu Lys Asn Gly Gly Pro Ser Ser Gly Ala Pro Pro Pro Ser
139
                  20
                                      25
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140
141
142
     (2) INFORMATION FOR SEQ ID NO:5:
143
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145
                (A) LENGTH: 39 amino acids
146
                (B) TYPE: amino acid
147
148
                (C) STRANDEDNESS: single
149
                (D) TOPOLOGY: linear
150
151
           (ii) MOLECULE TYPE: peptide
152
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(ix) FEATURE:
153
                (B) LOCATION: 39
154
                (D) OTHER INFORMATION: amidated Ser (Serineamide)
155
156
157
           (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
158
     His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
159
160
     Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
161
                                                            30
162
                                       25
                  20
163
     Ser Gly Ala Pro Pro Pro Ser
164
              35
165
     (2) INFORMATION FOR SEQ ID NO:6:
166
167
           (i) SEQUENCE CHARACTERISTICS:
168
                (A) LENGTH: 39 amino acids
169
170
                (B) TYPE: amino acid
                (C) STRANDEDNESS: single
171
172
                (D) TOPOLOGY: linear
173
         (ii) MOLECULE TYPE: peptide
174
175
176
         (ix) FEATURE:
                (B) LOCATION: 39
177
178
                (D) OTHER INFORMATION: amidated Ser (Serineamide)
179
180
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
181
     His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Leu Glu Glu
182
183
                                           10
     Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
184
185
                                       25
                                                            30
186
     Ser Gly Ala Pro Pro Pro Ser
187
188
189
190 (2) INFORMATION FOR SEQ ID NO:7:
191
192
          (i) SEQUENCE CHARACTERISTICS:
193
                (A) LENGTH: 39 amino acids
194
                (B) TYPE: amino acid
195
                (C) STRANDEDNESS: single
196
                (D) TOPOLOGY: linear
197
198
199
         (ii) MOLECULE TYPE: peptide
200
201
202
203
         (ix) FEATURE:
                (B) LOCATION: 39
204
                (D) OTHER INFORMATION: amidated Ser (Serineamide)
205
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206
207
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
208
209
210
     His Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
211
212
     Glu Ala Val Arg Leu Phe Ile Glu Phe Leu Lys Asn Gly Gly Pro Ser
                                                            30
213
                  20
                                       25
214
     Ser Gly Ala Pro Pro Pro Ser
215
              35
216
217
      (2) INFORMATION FOR SEQ ID NO:8:
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219
           (i) SEQUENCE CHARACTERISTICS:
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221
                (A) LENGTH: 39 amino acids
222
                (B) TYPE: amino acid
223
                (C) STRANDEDNESS: single
                (D) TOPOLOGY: linear
224
225
         (ii) MOLECULE TYPE: peptide
226
227
          (ix) FEATURE:
228
                (B) LOCATION: 39
229
                (D) OTHER INFORMATION: amidated Ser (Serineamide)
230
231
232
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
233
     Tyr Gly Glu Gly Thr Phe Thr Ser Asp Leu Ser Lys Gln Met Glu Glu
234
235
     Glu Ala Val Arg Leu Phe Ile Glu Trp Leu Lys Asn Gly Gly Pro Ser
236
237
                  20
                                       25
                                                            30
238
     Ser Gly Ala Pro Pro Pro Ser
239
              35
240
241
     (2) INFORMATION FOR SEQ ID NO:9:
242
243
           (i) SEQUENCE CHARACTERISTICS:
244
                (A) LENGTH: 39 amino acids
245
                (B) TYPE: amino acid
246
247
248
                (C) STRANDEDNESS: single
                (D) TOPOLOGY: linear
249
250
251
          (ii) MOLECULE TYPE: peptide
252
          (ix) FEATURE:
253
254
                (B) LOCATION: 39
                (D) OTHER INFORMATION: amidated Tyr (Tyrosinamide)
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256
257
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
258
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SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/08/908,867A*

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32 Wrong Classification (C) CLASSIFICATION: Pending 37 Wrong Classification (C) CLASSIFICATION: Pending